

## Sequence Listings

### (1) GENERAL INFORMATION:

- 5 (i) INVENTORS: B. Mollet  
R.D. Pridmore  
M.C. Zwahlen
- 10 (ii) TITLE OF THE INVENTION: Novel strains of the *Bacillus subtilis* group  
for food fermentation
- (iii) NUMBER OF SEQUENCES: 4
- 15 (iv) ADDRESS:  
(A) Address: Winston & Strawn  
(B) Street: 200 Park Avenue  
(C) City: New York  
(D) State: New York  
(E) Country: USA  
20 (F) Regional Code:
- (v) COMPUTER READABLE FORM:  
(A) Medium type: Floppy Disk  
(B) Computer: IBM PC Compatible  
25 (C) System: PC-DOS/MS-DOS  
(D) Software: Patentin Release Nr. 1, Version Nr. 1,25
- (vi) INFORMATION OF THE APPLICATION:  
(A) Application number: to be obtained  
30 (B) Application date: herewith  
(C) Classification:
- (vii) PRIORITY DATES: none
- 35 (viii) Attorney/Agent Information  
(A) Name: Allan A. Fanucci  
(B) Registration number: 30,256  
(C) Reference: 88265-4011
- 40 (ix) TELECOMMUNICATION:  
(A) Telephone: 212-294-3311  
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### (2) INFORMATION FOR SEQ ID. NO. 1:

- 45 (i) SEQUENCE CHARACTERISTICS:  
(A) length: 30 nucleotides  
(B) type: nucleic acid  
(C) strandedness: single stranded  
50 (D) topology: linear
- (ii) MOLECULE TYPE: DNA

(iii) SEQUENCE DESCRIPTION: SEQ ID. NO. 1:

GCGGCGGATC CGCTGATGAT CTCCCAGCCC

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5 (3) INFORMATION FOR SEQ ID. NO. 2:

(i) SEQUENCE CHARACTERISTICS:

(A) length: 44 nucleotides

(B) type: nucleic acid

(C) strandedness: single stranded

10 (D) topology: linear

(ii) MOLECULE TYPE: DNA

(iii) SEQUENCE DESCRIPTION: SEQ ID. NO. 2:

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CTCAAATTCC ATTCCTCAT CAGGACATGC ATAGCGTATC ATCC

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(4) INFORMATION FOR SEQ ID. NO. 3:

(i) SEQUENCE CHARACTERISTICS:

20 (A) length: 31 nucleotides

(B) type: nucleic acid

(C) strandedness: single stranded

(D) Topology: linear

25 (ii) MOLECULE TYPE: DNA

(iii) SEQUENCE DESCRIPTION: SEQ ID. NO. 3:

GGGGTCGAAT TCCACGAGAT ATCTAACTGC C

31

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(5) INFORMATION FOR SEQ ID. NO. 4:

(i) SEQUENCE CHARACTERISTICS:

(A) length: 44 nucleotides

(B) type: nucleic acid

35 (C) strandedness: single stranded

(D) Topology: linear

(ii) MOLECULE TYPE: DNA

40 (iii) SEQUENCE DESCRIPTION: SEQ ID. NO. 4:

GGATGATACG CTATGCATGT CCTGATGAGG AAATGGAATTTGAG

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